

S6 Table. Immunoprecipitation experiment analyzed by mass spectrometry.

Protein	MW (kDa)	Counts Input <i>ΔespL</i> /HA. <i>espL</i>	Counts IP H37Rv	Counts IP <i>ΔespL</i>	Counts IP <i>ΔespL/espL</i> .HA	Counts IP <i>ΔespL</i> /HA. <i>espL</i>
GroEL2	57	121	16	48	58	45
DnaK	67	19	8	8	13	11
<b>EspL.HA or HA.EspL</b>	<b>14</b>	<b>3</b>	<b>4</b>	<b>11</b>	<b>125</b>	<b>187</b>
RpoB	129	3	19	5	5	2
RpoC	147	4	4	0	18	2
SigA	58	3	27	11	16	14
Tuf	44	11	0	20	8	8
mIHF	21	10	0	9	3	0
GyrA	92	1	5	11	0	0
RplK	15	3	0	6	0	2
<b>EspD</b>	<b>20</b>	<b>0</b>	<b>5</b>	<b>0</b>	<b>11</b>	<b>35</b>
EsxB	11	5	0	0	5	0

The Table reports peptide counts for the indicated proteins in the samples obtained upon immunoprecipitation (IP) with anti-HA antibodies.

The total protein extract (Input) was used as a control. EspL.HA, HA.EspL and EspD are indicated in bold.

MW: molecular weight. H37Rv: wild type strain. *ΔespL*: *espL* knock-out mutant. *ΔespL/espL*.HA: *espL* knock-out mutant complemented by HA-tagged *espL* (HA at C-terminus). *ΔespL*/HA.*espL*: *espL* knock-out mutant complemented by HA-tagged *espL* (HA at N-terminus).